

REFERENCES

1. Reilly PA, Lehr T, Haertter S, et al. The effect of dabigatran plasma concentrations and patient characteristics on the frequency of ischemic stroke and major bleeding in atrial fibrillation patients: the RE-LY Trial (Randomized Evaluation of Long-Term Anticoagulation Therapy). *J Am Coll Cardiol* 2014;63:321-8.
2. Wessler JD, Grip LT, Mendell J, Giugliano RP. The P-glycoprotein transport system and cardiovascular drugs. *J Am Coll Cardiol* 2013;61:2495-502.
3. Ieiri I. Functional significance of genetic polymorphisms in P-glycoprotein (MDR1, ABCB1) and breast cancer resistance protein (BCRP, ABCG2). *Drug Metab Pharmacokinet* 2012;27:85-105.
4. McConeghy KW, Bress A, Qato DM, Wing C, Nutescu EA. Evaluation of dabigatran bleeding adverse reaction reports in the FDA Adverse Event Reporting System during the first year of approval. *Pharmacotherapy* 2014;34:561-9.

MicroRNA-29a Is a Friend or Foe for Cardiac Hypertrophy?



MicroRNAs (miRNAs) play an important role in the pathogenesis of structural alterations of cardiac hypertrophy (1). However, the precise mechanisms of miRNA-29a in cardiac hypertrophy are still unclear. A recent report in the *Journal* by Roncarati et al. (2) indicated that correlation with left ventricular hypertrophy parameters holds true for only miRNA-29a, which is significantly associated with both cardiac hypertrophy and fibrosis. This suggests that miRNA-29a is a potential therapeutic target for cardiac hypertrophy.

Other studies have found that down-regulation of miRNA-29a is associated with pathological cardiac hypertrophy. Abonnenc et al. (3) showed that levels of miRNA-29a were significantly reduced in a mouse model of pathological but not physiological hypertrophy (3). Furthermore, Divakaran et al. (4) indicated that Smad3 signaling plays dual roles in the heart: a beneficial role by delimiting hypertrophic growth and a deleterious role by modulating myocardial fibrosis, possibly through a pathway that entails accumulation of miRNA-29a, which decreases collagen gene expression. However, Zile et al. (5) showed that miRNA-29a levels increased 5 days after myocardial infarction and then decreased to control levels at later time points. A time-dependent change in miRNA-29a levels occurred in patients with myocardial infarction, including an early and robust increase in miRNA levels that affected myocardial fibrosis.

These findings suggest that miRNA-29a may be a good clinical biomarker for patients with cardiac hypertrophy because it seems to play a key role in the development of this condition. Additional studies

with more patients and animal studies are needed to determine the precise role of miRNA-29a in cardiac hypertrophy, and therapeutic agents targeting miRNA-29a might result in innovative new therapies. All in all, we greatly enjoyed reading the article by Roncarati et al. (2) and believe that miRNA-29a may be useful for the prevention and treatment of cardiac hypertrophy.

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REFERENCES

1. Mathiyalagan P, Okabe J, Chang L, Su Y, Du XJ, El-Osta A. The primary microRNA-208b interacts with Polycomb-group protein, Ezh2, to regulate gene expression in the heart. *Nucleic Acids Res* 2014;42:790-803.
2. Roncarati R, Viviani Anselmi C, Losi MA, et al. Circulating miR-29a, among other up-regulated microRNAs, is the only biomarker for both hypertrophy and fibrosis in patients with hypertrophic cardiomyopathy. *J Am Coll Cardiol* 2014;63:920-7.
3. Abonnenc M, Nabeebaccus AA, Mayr U, et al. Extracellular matrix secretion by cardiac fibroblasts: role of microRNA-29b and microRNA-30c. *Circ Res* 2013;113:1138-47.
4. Divakaran V, Adroque J, Ishiyama M, et al. Adaptive and maladaptive effects of SMAD3 signaling in the adult heart after hemodynamic pressure overloading. *Circ Heart Fail* 2009;2:633-42.
5. Zile MR, Mehurg SM, Arroyo JE, Stroud RE, DeSantis SM, Spinale FG. Relationship between the temporal profile of plasma microRNA and left ventricular remodeling in patients after myocardial infarction. *Circ Cardiovasc Genet* 2011;4:614-9.

Pro-Inflammatory Interleukin Genotypes Potentiate Early and Advanced Atherosclerosis Differently



We read with interest the paper by Tsimikas et al. (1) in a recent issue of the *Journal*, which reported that genetic differences in the interleukin (IL)-1 gene cluster, known to be associated with inflammatory responsiveness, strongly influence the presence of